

Task 1. Data explorations (30%)

Dog breeds

studentNumber

The dataset for this task contains information on different dog breeds (`dog_breed`). The variables that you require for this assignment are:

- `category`: breed category
- `lifetime_costs`: expected lifetime costs in dollars
- `suitability_for_children`: suitability for children (1 = high, 2 = medium, 3 = low)
- `longevity`: mean life expectancy in years
- `popularity_inUS_ranking`: popularity in the US, 1 = most popular, higher values indicates lower popularity

```
# Load packages

library(tidyverse)
library(foreign)

# Import data:
dogbreeds <- read_csv2("DogBreeds_selected.csv")
```

Template file and submission:

Add your code to the provided template file. Write reproducible and readable code to make the plots for both exercise a and b and make sure that the plots are visible in your output file (.html/.pdf). Keep the data stored in the subfolder so your .Rmd file can reach it.

For more submission instructions, please see the general instructions of this Graded Assignment.

a) Data descriptives - distributions per variable

Create suitable data visualizations to show the individual distributions of the breeds' mean life expectancy (`longevity`), suitability for children (`suitability_for_children`), categories (`category`) and expected lifetime costs (`lifetime_costs`). For each plot, only use the data that is available, i.e. exclude missing observations.

```
library(RColorBrewer)
library(ggmosaic)
library(see)
```

```
summary(dogbreeds)
```

##	dog_breed	category	datadog_score	popularity_inUS_ranking
##	Length:174	Length:174	Min. :0.990	Min. : 1.0
##	Class :character	Class :character	1st Qu.:2.185	1st Qu.:22.5
##	Mode :character	Mode :character	Median :2.710	Median :44.0
##			Mean :2.604	Mean :44.0

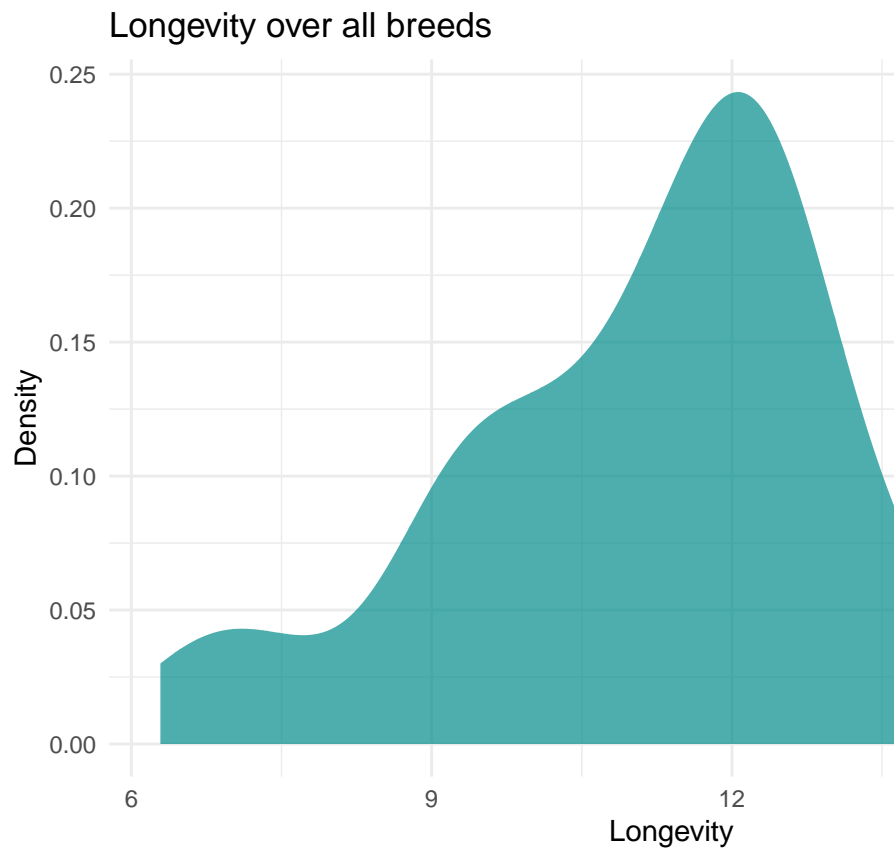
```
##              3rd Qu.:3.035    3rd Qu.:65.5
##              Max.      :3.640    Max.      :87.0
##              NA's      :87      NA's      :87
##   longevity grooming_required suitability_for_children size_category
##   Min.      : 6.29    Min.      :1.000    Min.      :1.000    Length:174
##   1st Qu.: 9.70    1st Qu.:2.000    1st Qu.:1.000    Class :character
##   Median :11.29    Median :2.000    Median :1.000    Mode  :character
##   Mean   :10.96    Mean   :1.804    Mean   :1.491
##   3rd Qu.:12.37    3rd Qu.:2.000    3rd Qu.:2.000
##   Max.   :16.50    Max.   :3.000    Max.   :3.000
##   NA's   :39      NA's   :62      NA's   :62
## intelligence_category lifetime_costs average_purchase_price
## Length:174           Min.      :11100    Min.      : 283.0
## Class :character     1st Qu.:15386    1st Qu.: 587.2
## Mode  :character     Median :17336    Median : 795.0
##                      Mean   :17069    Mean   : 876.8
##                      3rd Qu.:18861    3rd Qu.:1042.2
##                      Max.   :22640    Max.   :3460.0
##                      NA's   :83      NA's   :28
## price_category
## Length:174
## Class :character
## Mode  :character
##
##
##
```

```
cleaned <- na.omit(dogbreeds)
```

Note: These plots are expected to be self-contained (i.e. readers should be able to understand them without extra explanation) and to obey the principles of good graphics, but they are not meant to be formal presentation graphics. For example, you are not expected to use additional information to make the plot information rich. The focus is on uncovering the distributions of the variables.

```
# your code here
```

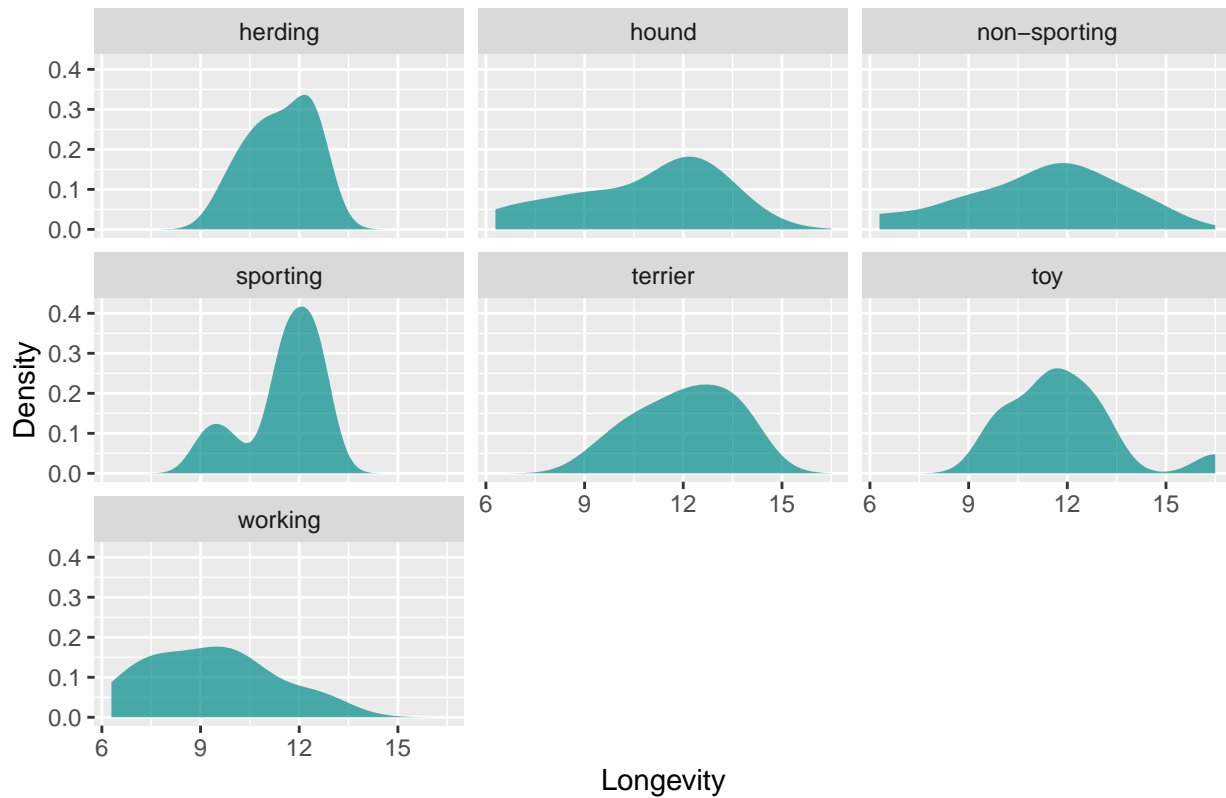
```
ggplot(data = cleaned, aes(x = longevity)) + geom_density(fill="darkcyan",
                                                           alpha = 0.7,
                                                           color=NA) +
  labs(x= "Longevity", y = "Density") + ggtitle("Longevity over all breeds") + theme_minimal()
```



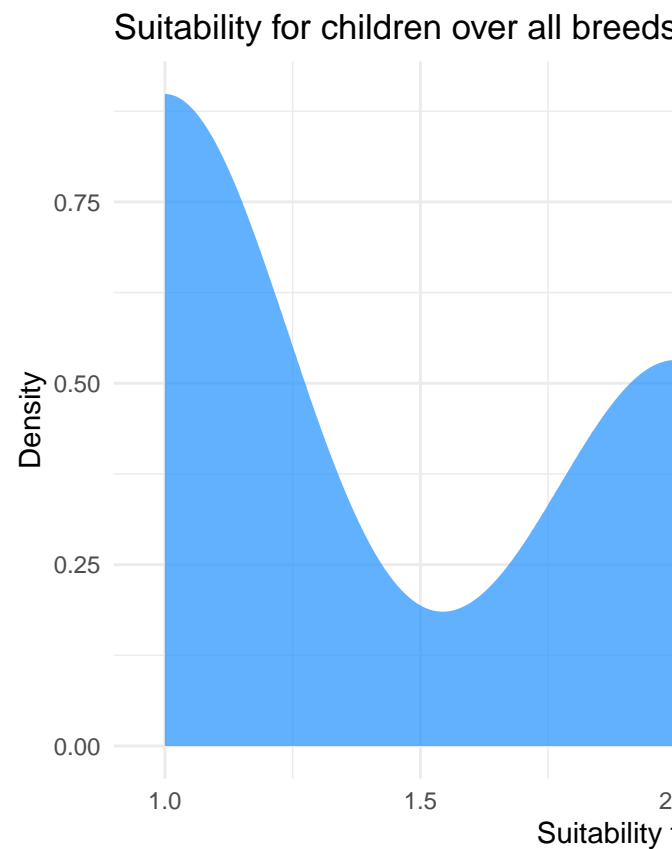
Explore mean life expectancy (longevity)

```
ggplot(data = cleaned, aes(x = longevity)) + geom_density(fill="darkcyan",  
                                                         alpha = 0.7,  
                                                         color=NA) +  
labs(x= "Longevity", y = "Density") + ggtitle("Longevity over all breeds for specific categories") + fa
```

Longevity over all breeds for specific categories



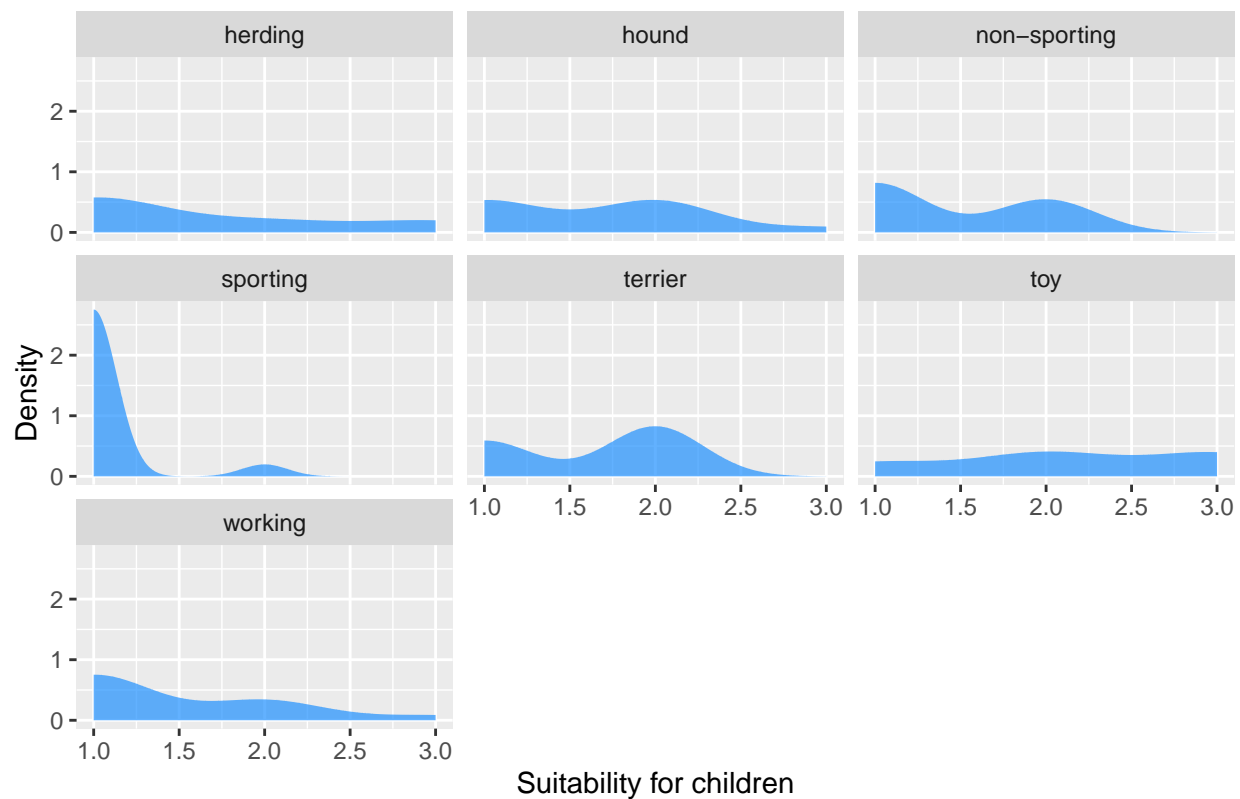
```
# your code here
ggplot(data = cleaned, aes(x = suitability_for_children)) + geom_density(fill="dodgerblue",
                                                                    alpha = 0.7,
                                                                    color=NA) +
  labs(x= "Suitability for children", y ="Density") + ggtitle("Suitability for children over all breeds")
```



Explore suitability for children (suitability_for_children)

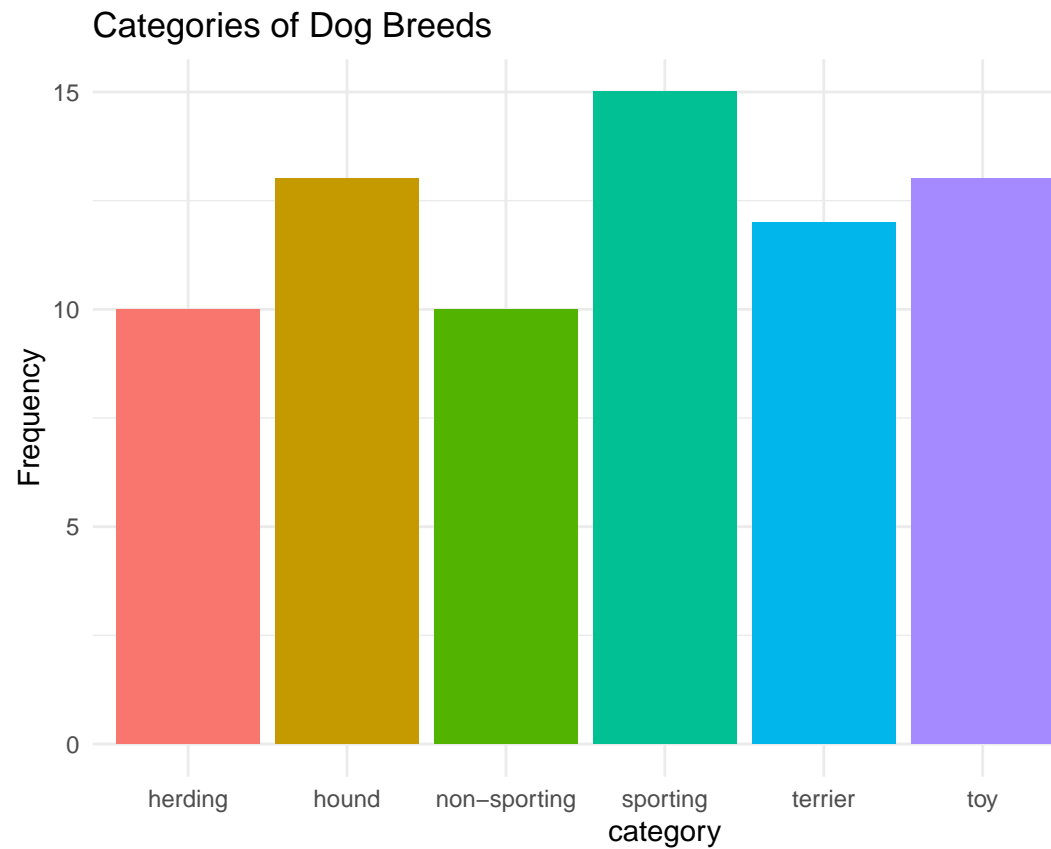
```
ggplot(data = cleaned, aes(x = suitability_for_children)) + geom_density(fill="dodgerblue",  
                                                                    alpha = 0.7,  
                                                                    color=NA) +  
  labs(x= "Suitability for children", y ="Density") + ggtitle("Suitability for children in specific categories")
```

Suitability for children in specific categories



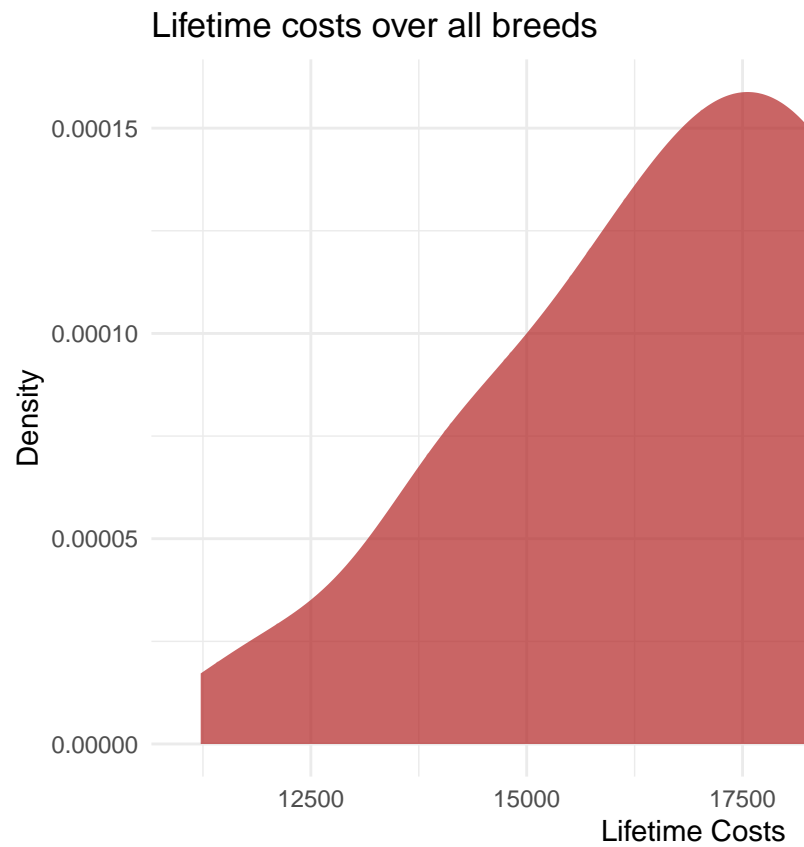
your code here

```
ggplot(data = cleaned,
       mapping = aes(
         x = category,
         fill = category
       )) + geom_bar(stat="count") + theme_minimal() + theme(legend.position="None") + ggtitle("Categories")
```



Explore categories (category)

```
ggplot(data = cleaned, aes(x = lifetime_costs)) + geom_density(fill="firebrick",alpha = 0.7,  
                                                                  color=NA) +  
  labs(x= "Lifetime Costs", y ="Density") +ggtitle("Lifetime costs over all breeds") + theme_minimal()
```

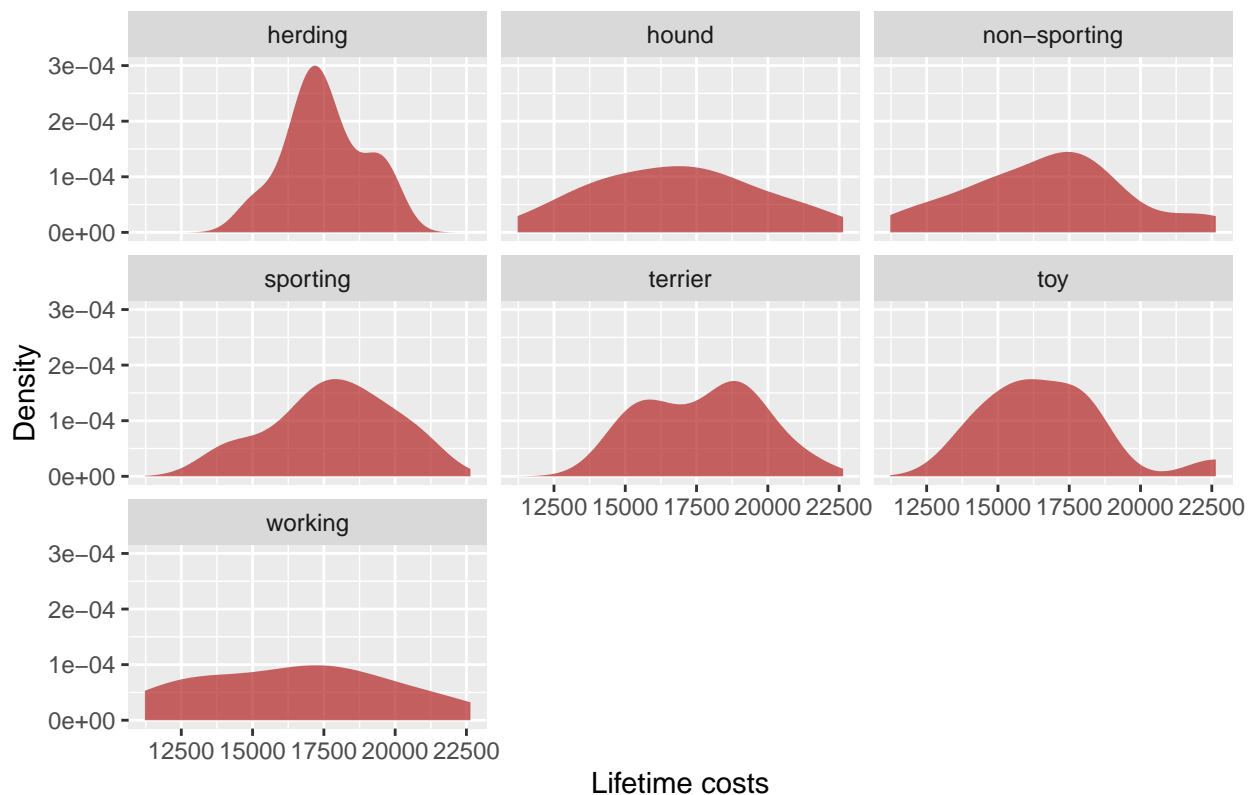


Explore expected lifetime costs (lifetime_costs)

your code here

```
ggplot(data = cleaned, aes(x = lifetime_costs)) + geom_density(fill="firebrick",  
                                                                alpha = 0.7,  
                                                                color=NA) +  
  labs(x= "Lifetime costs", y = "Density") + ggtitle("Lifetime costs in specific categories") + facet_wrap
```


Lifetime costs in specific categories



b) Data descriptives - relationships among 3 variables

Assume that you are studying reasons why some dog breeds are more popular than others. Your main hypothesis is that a dog breed's popularity in the US (`popularity_inUS_ranking`) depends on the expected lifetime costs of the breed (`lifetime_costs`). Additionally, you think that a breed's suitability for children (`suitability_for_children`) might be relevant.

Make a visualisation that focuses mostly on exploring your main hypothesis that popularity depends on lifetime costs, but also gives an indication of whether the breed's suitability for children has an influence.

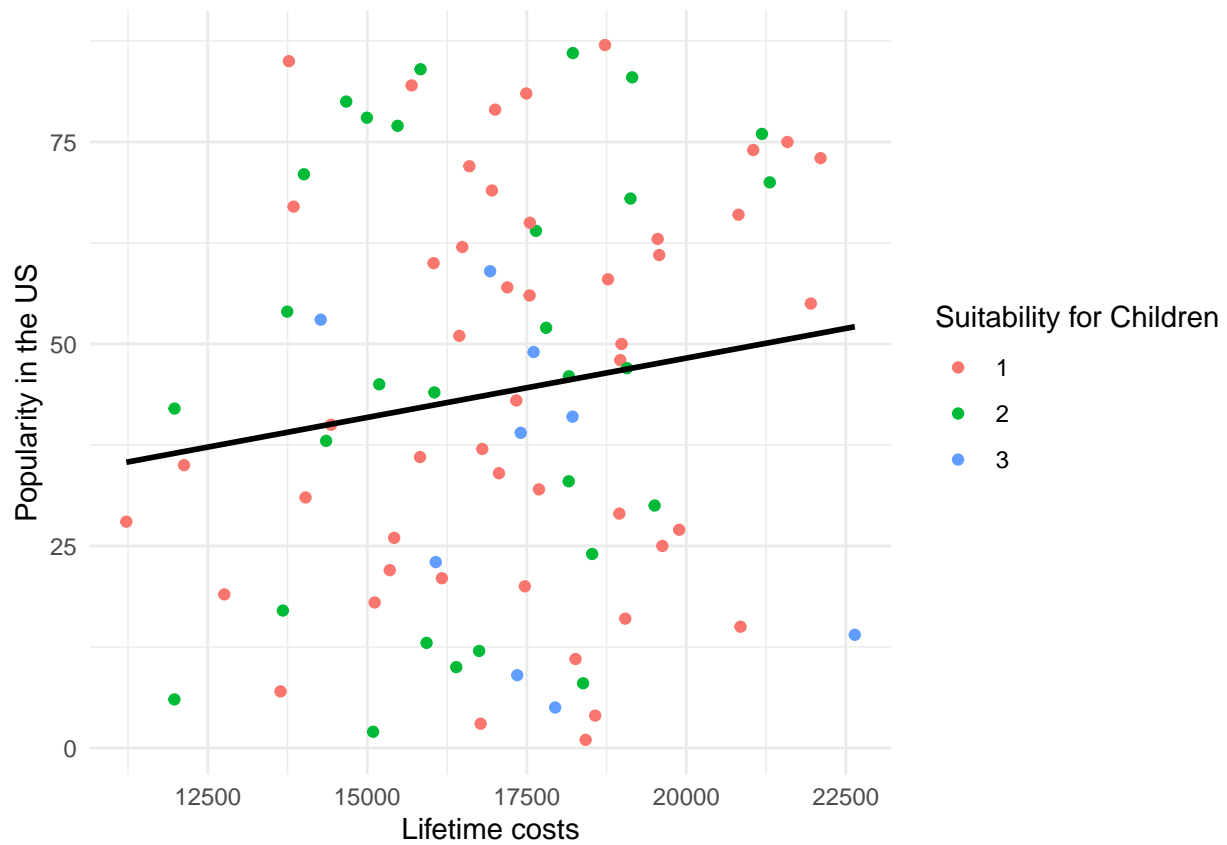
Note: This plot is expected to be self-contained (i.e. readers should be able to understand them without extra explanation) and to obey the principles of good graphics, but it is not meant to be formal presentation graphics. For example, you are not expected to use additional information to make the plot information rich. The focus is on uncovering the relationship among the variables.

your code here

```
ggplot(data = cleaned,
  mapping = aes(
    x = lifetime_costs,
    y = popularity_inUS_ranking

  )) + geom_point( aes(color = as.factor(suitability_for_children))) +
  geom_smooth(method="lm", se=FALSE, color="black") + theme_minimal() +
  labs(x="Lifetime costs", y="Popularity in the US", color="Suitability for Children")
```

```
## `geom_smooth()` using formula = 'y ~ x'
```



From the graph above it is safe to say that while there is a positive relationship between lifetime costs and popularity in the US, there is very little effect of suitability for children on popularity (seen by the points both above and below).